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- (1) nucleotide 5573 wherein X is T;
- (m) nucleotide 5659 wherein X is C;
- (n) nucleotide 5678 wherein X is C;
- (o) nucleotide 5874 wherein X is T; and
- (p) nucleotide 5934 wherein X is G

or the complement thereof.

- 2. 18 The isolated nucleix acid probe of claim & comprising at least two of:
- (a) nucleotide 194 wherein X is G;
- (b) nucleotide 294 wherein X\is T;
- (c) nucleotide 1136 wherein X is A;
- (d) nucleotide 1252 wherein X is T;
- (e) nucleotide 1334 wherein X is A;
- (f) nucleotide 1699 wherein X is C
- (g) nucleotide 3150 wherein X is G
- (h) nucleotide 3207 wherein X is T;
- (i) nucleotide 3209 wherein X is A;
- (i) nucleotide 5444 wherein X is C;
- (k) nucleotide 5551 wherein X is A;
- (1) nucleotide 5573 wherein X is T;
- (m) nucleotide 5659 wherein X is C;
- (n) nucleotide 5678 wherein X is C;
- (o) nucleotide 5874 wherein X is T; and
- (p) nucleotide 5934 wherein X is G

or the complement thereof.

2. 19 The probe of claim 1/comprising no more than 500 contiguous nucleotides of SEQ ID NO:1.

A. 20 The probe of claim 1 comprising no more than 200 contiguous nucleotides of SEQ ID NO:1.

5.21 The probe of claim 1/comprising no more than 100 contiguous nucleotides of SEO ID NO:1.

6.22 The probe of claim / comprising no more than 50 contiguous nucleotides of SEQ ID NO:1

7.23 The probe of claim 2 comprising DNA.

8/24 The probe of claim 1 comprising a peptide nucleic acid.

9.25 The probe of claim, further comprising a detectable label.



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The probe of claim wherein the detectable label is a fluorescent label.

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11.27 A method comprising

- (a) providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;
- (b) contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:1, the probe comprising at least one of:
 - (i) nucleotide 194 wherein X is G;
 - (ii) nucleotide 294 wherein X is\T;
 - (iii) nucleotide 1136 wherein X is A;
 - (iv) nucleotide 1252 wherein X is T;
 - (v) nucleotide 1334 wherein X is A;
 - (vi) nucleotide 1699 wherein X is C
 - (vii) nucleotide 3150 wherein X is G:
 - (viii) nucleotide 3207 wherein X is T;
 - (ixi) nucleotide 3209 wherein X is A
 - (x) nucleotide 5444 wherein X is C;
 - (xi) nucleotide 5551 wherein X is A;
 - (xii) nucleotide 5573 wherein X is T;
 - (xiii) nucleotide 5659 wherein X is k
 - (xiv) nucleotide 5678 wherein X is C:
 - (xv) nucleotide 5874 wherein X is T; and
 - (xvi) nucleotide 5934 wherein X is G

or the complement thereof; and

- (c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.
- 12.23 An isolated nucleic acid probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:2, the probe comprising at least one of:
 - (a) nucleotide 3434 wherein X is T;
 - (b) nucleotide 4313 wherein X is C;
 - (c) nucleotide 4799 wherein X is G;
 - (d) nucleotide 5255 wherein X is T;
 - (e) nucleotide 5455 wherein X is A;
 - (f) nucleotide 5507 wherein X is C:
 - (g) nucleotide 5810 wherein X is T;
 - (h) nucleotide 6128 wherein X is T;
 - (i) nucleotide 6626 wherein X is T;
- (j) nucleotide 6686 wherein X is T; or the complement thereof.
 - 13.29 The isolated nucleic acid probe of claim 12 comprising at least two of:
 - (a) nucleotide 3434 wherein X is T;

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(b) nucleotide 4313 wherein X is C;

- (c) nucleotide 4799 wherein X is G;
- (d) nucleotide 5255 wherein X is T;
- (e) nucleotide 5455 wherein X is A;
- (f) nucleotide 5507 wherein X is C;
- (g) nucleotide 5810 wherein X is T;
- (h) nucleotide 6128 wherein X is T;
- (i) nucleotide 6626 wherein X is T:
- (j) nucleotide 6686 wherein X\is T; or the complement thereof.

14.30 The probe of claim 12 comprising no more than 500 contiguous nucleotides of SEQ ID NO:1.

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15.3 The probe of claim 12 comprising no more than 200 contiguous nucleotides of SEO ID NO:2.

16.32 The probe of claim 12 comprising no more than 100 contiguous nucleotides of SEQ ID NO:2.

The probe of claim 12 comprising no more than 50 contiguous nucleotides of SEQ ID NO:2.

18.34 The probe of claim 12 comprising DNA

19.35 The probe of claim 12 comprising a peptide nucleic acid.

20.36 The probe of claim 12 further comprising a detectable label.

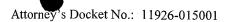
21.37 The probe of claim 20 wherein the detectable label is a fluorescent label.

22.38 A method comprising:

(a) providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;

- (b) contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:2, the probe comprising at least one of:
 - (i) nucleotide 3434 wherein X is T;
 - (ii) nucleotide 4313 wherein X is C;
 - (iii) nucleotide 4799 wherein X is G;
 - (iv) nucleotide 5255 wherein X is T;
 - (v) nucleotide 5455 wherein X is A;
 - (vi) nucleotide 5507 wherein X is C;
 - (vii) nucleotide 5810 wherein X is T;
 - (viii) nucleotide 6128 wherein X is T;

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- (ixi) nucleotide 6626 wherein X is T;
- (x) nucleotide 6686 wherein X is T;

or the complement thereof; and

- (c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.
- 23. ³⁹ An isolated nucleic acid probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:3, the probe comprising at least one of:
 - (a) nucleotide 166 wherein X is C;
 - (b) nucleotide 577 wherein X is G;
 - (c) nucleotide 638 wherein X is G;
 - (d) nucleotide 1708 wherein X is G;
 - (e) nucleotide 3432 wherein X is C;
 - (f) nucleotide 3682 wherein X is T;
 - (g) nucleotide 3730 wherein X is A;
 - (b) medicatide 2025 wherein Win C
 - (h) nucleotide 3925 wherein X\is G;
- (i) nucleotide 3937 wherein X is C; or the complement thereof.

24.40 The isolated nucleic acid probe of claim 23 comprising at least two of:

- (a) nucleotide 166 wherein X s C;
- (b) nucleotide 577 wherein X is G;
- (c) nucleotide 638 wherein X is G
- (d) nucleotide 1708 wherein X is G;
- (e) nucleotide 3432 wherein \hat{X} is C;
- (f) nucleotide 3682 wherein X is T;
- (g) nucleotide 3730 wherein X is A;
- (h) nucleotide 3925 wherein X is G;
- (i) nucleotide 3937 wherein X is C; or the complement thereof.

25.41 The probe of claim 23 comprising no more than 500 contiguous nucleotides of SEQ ID NO:3.

26.42 The probe of claim 23 comprising no more than 200 contiguous nucleotides of SEQ ID NO:3.

21.43 The probe of claim 23 comprising no more than 100 contiguous nucleotides of SEQ ID NO:3.

28. The probe of claim 23 comprising no more than 50 contiguous nucleotides of SEQ ID NO:3.

29.45 The probe of claim 23 comprising DNA.

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36. 46 The probe of claim 23 comprising a peptide nucleic acid.

21.47 The probe of claim 23 further comprising a detectable label.

32.44 The probe of claim, 34 wherein the detectable label is a fluorescent label.

33.49 A method comprising:

providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;

- contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:3, the probe comprising at least one of:
 - (i) nucleotide 166 wherein X is C:
 - (ii) nucleotide 577 wherein X is G;
 - (iii) nucleotide 638 wherein X is G;
 - (iv) nucleotide 1708 wherein X is G;
 - (v) nucleotide 3432 wherein X is C:
 - (vi) nucleotide 3682 wherein X is\T;
 - (vii) nucleotide 3730 wherein X is A:
 - (viii) nucleotide 3925 wherein X is
 - (ixi) nucleotide 3937 wherein X is

or the complement thereof; and

- (c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.
- 34.50 An isolated nucleic acid probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:4, the probe comprising at least one of:
 - (a) nucleotide 175 wherein X is A;
 - (b) nucleotide 341 wherein X is G;
 - (c) nucleotide 791 wherein X is T;
 - (d) nucleotide 1067 wherein X is A;
 - (e) nucleotide 1337 wherein X is A:
 - (f) nucleotide 1997 wherein X is C:
 - (g) G is inserted after nucleotide 2106;
 - (h) nucleotide 2582 wherein X is G;
 - (i) nucleotide 2617 wherein X is T;
 - (j) nucleotide 2652 wherein X is C;

or the complement thereof.

- 25.5 The isolated nucleic acid probe of claim 34 comprising at least two of:
- (a) nucleotide 175 wherein X is A:
- (b) nucleotide 341 wherein X is G:
- (c) nucleotide 791 wherein X is T;
- (d) nucleotide 1067 wherein X is A;

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(e) nucleotide 1337 wherein X is A;

- (f) nucleotide 1997 wherein X is C;
- (g) G is inserted after nucleotide 2100;
- (h) nucleotide 2582 wherein X is G;
- (i) nucleotide 2617 wherein X is T;
- (j) nucleotide 2652 wherein X is C; or the complement thereof.

36. 52 The probe of claim 34 comprising no more than 500 contiguous nucleotides of SEQ ID NO:4.

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37.53 The probe of claim 34 comprising no more than 200 contiguous nucleotides of SEQ ID NO:4.

38.54 The probe of claim 34 comprising no more than 100 contiguous nucleotides of SEO ID NO:4.

39.55 The probe of claim 34 comprising no-more than 50 contiguous nucleotides of SEQ ID NO:4.

40.5 6The probe of claim 34 comprising DNA.

A1.57 The probe of claim 34 comprising a peptide nucleic acid.

42.5 The probe of claim 44 further comprising a detectable label.

43.59 The probe of claim 42 wherein the detectable label is a fluorescent label.

44.60 A method comprising:

- (a) providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;
- (b) contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:4, the probe comprising at least one of:
 - (i) nucleotide 175 wherein X is A;
 - (ii) nucleotide 341 wherein X is G;
 - (iii) nucleotide 791 wherein X is T;
 - (iv) nucleotide 1067 wherein X is A
 - (v) nucleotide 1337 wherein X is A;
 - (vi) nucleotide 1997 wherein X is C;
 - (vii) wherein G is inserted after nuclebtide 2100:
 - (viii) nucleotide 2582 wherein X is G
 - (ixi) nucleotide 2617 wherein X is T;
- (x) nucleotide 2652 wherein X is C; or the complement thereof; and

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(c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.

A5.4 An isolated nucleic acid probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO 5, the probe comprising at least one of:

- (a) nucleotide 431 wherein X is G:
- (b) nucleotide 441 wherein X is G;
- (c) nucleotide 498 wherein X is Γ ;
- (d) nucleotide 579 wherein X is C;
- (e) nucleotide 599 wherein X is C; or the complement thereof.

46.62 The isolated nucleic acid probe of claim 45 comprising at least two of:

- (a) nucleotide 431 wherein X is G;
- (b) nucleotide 441 wherein X is C
- (c) nucleotide 498 wherein X is T
- (d) nucleotide 579 wherein X is C
- (e) nucleotide 599 wherein X is C or the complement thereof.

AT 63 The probe of claim 45 comprising no more than 500 contiguous nucleotides of

4864 The probe of claim 43 comprising no more than 200 contiguous nucleotides of

4965 The probe of claim 45 comprising no more than 100 contiguous nucleotides of

50.66 The probe of claim 45 comprising no more than 50 contiguous nucleotides of SEO ID NO:5.

81.67 The probe claim 48 comprising DNA.

5268 The probe of claim 45 comprising a peptide nucleic acid.

5369 The probe of claim 45 further comprising a detectable label.

54.70 The probe of claim 53 wherein the detectable label is a fluorescent label.

55.7/ A method comprising:

providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;

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(b) contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:5, the probe comprising at least one of:

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- (i) nucleotide 431 wherein X is G;
- (ii) nucleotide 441 wherein X is G;
- (iii) nucleotide 498 wherein X is T;
- (iv) nucleotide 579 wherein X is C;
- (v) nucleotide 599 wherein X is C;

or the complement thereof; and

(c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.

56.72 An isolated nucleic acid probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:6, the probe comprising at least one of:

- (a) nucleotide 1066 wherein X is C;
- (b) nucleotide 1136 wherein X is G;
- (c) nucleotide 1497 wherein X is A; or the complement thereof.

51.73 The isolated nucleic acid probe of claim 56 comprising at least two of:

- (a) nucleotide 1066 wherein X is C:
- (b) nucleotide 1136 wherein X is G;
- (c) nucleotide 1497 wherein X is A; or the complement thereof.

58 74 The probe of claim 56 comprising no more than 500 contiguous nucleotides of SEQ ID NO:6.

59.75 The probe of claim 56 comprising no more than 200 contiguous nucleotides of SEQ ID NO:6.

60. 76 The probe of claim 56 comprising no more than 100 contiguous nucleotides of SEQ ID NO:6.

61.77 The probe of claim 56 comprising no more than 50 contiguous nucleotides of SEQ ID NO:6.

62.78 The probe of claim 56 comprising DNA.

63.79 The probe of claim 56 comprising a pentide nucleic acid.

64.80 The probe of claim 36 further comprising a detectable label.

65.81 The probe of claim 64 wherein the detectable label is a fluorescent label.

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66.82 A method comprising:

(a) providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;

(b) contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:6, the probe comprising at least one of:

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- (i) nucleotide 1066 wherein X is C;
- (ii) nucleotide 1136 wherein X is G;
- (iii) nucleotide 1497 wherein X is A;

or the complement thereof; and

(c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.

67. 1/3 An isolated nucleic acid probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:7, the probe comprising at least one of:

- (a) nucleotide 276 wherein X is T;
- (b) nucleotide 321 wherein X is C;
- (c) nucleotide 452 wherein X is A;
- (d) C is inserted after nucleotide 45%;
- (e) nucleotide 491 wherein X is A;
- (f) nucleotide 533 wherein X is C;
- (g) nucleotide 624 wherein X is C;
- (h) nucleotide 639 wherein X is G;
- (i) nucleotide 655 wherein X is C; or the complement thereof.

68. 84 The isolated nucleic acid probe of claim 67 comprising at least two of:

- (a) nucleotide 276 wherein X is T;
- (b) nucleotide 321 wherein X is C;
- (c) nucleotide 452 wherein X is A;
- (d) C is inserted after nucleotide 457;
- (e) nucleotide 491 wherein X is A;
- (f) nucleotide 533 wherein X is C;
- (g) nucleotide 624 wherein X is C;
- (h) nucleotide 639 wherein X is G;
- (i) nucleotide 655 wherein X is C;

or the complement thereof.

69.85 The probe of claim 67 comprising no more than 500 contiguous nucleotides of SEQ ID NO:7.

70.86 The probe of claim 67 comprising no more than 200 contiguous nucleotides of SEQ ID NO:7.

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71.87 The probe of claim 67 comprising no more than 100 contiguous nucleotides of SEQ ID NO:7.

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72.88 The probe of claim 67 comprising no more than 50 contiguous nucleotides of SEQ ID NO:7.

73.89 The probe of claim 67 comprising DNA.

74.90 The probe of claim 67 comprising a peptide nucleic acid.

75.9 The probe of claim of further comprising a detectable label.

76.92 The probe of claim I wherein the detectable label is a fluorescent label.

71.93 A method comprising:

(a) providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;

(b) contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQID NO:7, the probe comprising at least one of:

(i) nucleotide 276 wherein X is T

(ii) nucleotide 321 wherein X is C

(iii) nucleotide 452 wherein X is A;

(iv) C is inserted after nucleotide 457;

(v) nucleotide 491 wherein X is A;

(vi) nucleotide 533 wherein X is

(vii) nucleotide 624 wherein X is C; (viii) nucleotide 639 wherein X is G;

(ixi) nucleotide 655 wherein X is C;

or the complement thereof; and

(c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.

78. 94 An isolated nucleic acid probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:8, the probe comprising at least one of:

(a) nucleotide 701 wherein X is ¢;

(b) nucleotide 716 wherein X is G;

(c) nucleotide 732 wherein X is C

(d) nucleotide 1293 wherein X is G;

(e) nucleotide 1322 wherein X is G;

(f) nucleotide 1379 wherein X is C

(g) nucleotide 1590 wherein X is T

(h) nucleotide 1688 wherein X is G; (i) nucleotide 2401 wherein X is G;

(i) nucleotide 2429 wherein X is A

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(k) nucleotide 2488 wherein X is T;

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(1) nucleotide 2594 wherein X is T;

(m) nucleotide 2618 wherein X is A;

(n) nucleotide 3083 wherein X is A;

(o) nucleotide 3125 wherein X is A;

(p) nucleotide 3212 wherein X is T;

(q) nucleotide 3619 wherein X is A;

(r) nucleotide 3635 wherein X is A;

(s) nucleotide 4256 wherein X is A;

(t) nucleotide 4898 wherein X is G;

(u) nucleotide 5006 wherein X is T;

(v) nucleotide 5062 wherein X is A:

(w) nucleotide 5167 wherein X is A;

(x) nucleotide 11069 wherein X is G;

(y) nucleotide 11238 wherein X is T;

(z) nucleotide 11293 wherein X is G:

(aa) nucleotide 11422 wherein X is C;

(bb) nucleotide 11686 wherein X is T;

(cc) nucleotide 12598 wherein X is C;

(dd) nucleotide 13171 wherein X is C;

(ee) nucleotide 13298 wherein X is A;

(ff) nucleotide 13645 wherein X is C;

(gg) nucleotide 13751 wherein X is A

(hh) nucleotide 13782 wherein X is C;

(ii) nucleotide 13806 wherein X/s C;

(jj) nucleotide 13813 wherein X is C;

(kk) nucleotide 14479 wherein X is G;

(ll) T is inserted after nucleotide 14546;

(mm) nucleotide 14585 wherein X is T;

(nn) nucleotide 14729 wherein X is A;

(oo) nucleotide 14787 wherein X is T:

(pp) nucleotide 14795 whereih X is A;

(qq) nucleotide 15041 wherein X is C;

(rr) nucleotide 15343 wherein X is A;

(ss) nucleotide 15449 wherein X is A:

(tt) nucleotide 15502 wherein X is A;

(uu) nucleotide 15545 wherein X is T:

(vv) nucleotide 15589 wherein\X is G;

(ww) nucleotide 15769 wherein X is T;

(xx) nucleotide 15839 wherein X is G;

(yy) nucleotide 16148 wherein X is A:

(zz) nucleotide 16198 wherein X is G; and

(aaa) nucleotide 16202 wherein X is T;

or the complement thereof.

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79. 95 The isolated nucleic acid probe of claim 78 comprising at least two of:

(a) nucleotide 701 wherein X is C;

(b) nucleotide 716 wherein X is G;

(c) nucleotide 732 wherein X is C;

(d) nucleotide 1293 wherein X is G;

(e) nucleotide 1322 wherein X is G;

(f) nucleotide 1379 wherein X is C;

(g) nucleotide 1590 wherein X is T;

(h) nucleotide 1688 wherein X is G;

(i) nucleotide 2401 wherein X is G;

(j) nucleotide 2429 wherein X is A;

(k) nucleotide 2488 wherein X is T;

(1) nucleotide 2594 wherein X is T;

(m) nucleotide 2618 wherein X is A;

(n) nucleotide 3083 wherein X is A;

(o) nucleotide 3125 wherein X is A;

(p) nucleotide 3212 wherein X is T;

(q) nucleotide 3619 wherein X is A;

(r) nucleotide 3635 wherein X is A;

(s) nucleotide 4256 wherein X is A;

(t) nucleotide 4898 wherein X is G;

(u) nucleotide 5006 wherein X is T;

(v) nucleotide 5062 wherein X is A;

(w) nucleotide 5167 wherein X is A;

(x) nucleotide 11069 wherein X is G;

(y) nucleotide 11238 wherein X is T; (z) nucleotide 11293 wherein X is G;

(aa) nucleotide 11422 wherein X is C;

(bb) nucleotide 11686 wherein X is T;

(cc) nucleotide 12598 wherein X is C;

(cc) nucleotide 12598 wherein X is C; (dd) nucleotide 13171 wherein X is C;

(ee) nucleotide 13298 wherein X is A;

(ff) nucleotide 13645 wherein X is C;

(gg) nucleotide 13751 wherein X is A;

(hh) nucleotide 13782 wherein X is C;

(ii) nucleotide 13806 wherein X is C;

(jj) nucleotide 13813 wherein X is C;

(kk) nucleotide 14479 wherein X is G;

(ll) nucleotide 14546 wherein X\is T;

(mm) T is inserted after nucleotide 14546;

(nn) nucleotide 14729 wherein X is A;

(00) nucleotide 14787 wherein X is T;

(pp) nucleotide 14795 wherein X is A;

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(qq) nucleotide 15041 wherein X is C;

(rr) nucleotide 15343 wherein X is A;

(ss) nucleotide 15449 wherein X is A;

(tt) nucleotide 15502 wherein X is A;

(uu) nucleotide 15545 wherein X is T;

(vv) nucleotide 15589 wherein X is G;

(ww) nucleotide 15769 wherein X is T;

(xx) nucleotide 15839 wherein X is G;

(yy) nucleotide 16148 wherein X is A;

(zz) nucleotide 16198 wherein X is G; and

(aaa) nucleotide 16202 wherein X is T;

or the complement thereof.

80. The probe of claim 78 comprising no more than 500 contiguous nucleotides of SEQ ID NO:8.

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81.97 The probe of claim 78 comprising no more than 200 contiguous nucleotides of SEQ ID NO:8.

82. 9 The probe of claim 78 comprising no more than 100 contiguous nucleotides of SEQ ID NO:8.

83.99 The probe of claim 78 comprising no more than 50 contiguous nucleotides of SEQ ID NO:8.

84. // The probe of claim \$ comprising DNA.

85./0/ The probe of claim 78 comprising a peptide nucleic acid.

86./02 The probe of claim 18 further comprising a detectable label.

87./03 The probe of claim 86 wherein the detectable label is a fluorescent label.

88.104 A method comprising:

- (a) providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;
- (b) contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:8, the probe comprising at least one of:
 - (i) nucleotide 701 wherein X is C;
 - (ii) nucleotide 716 wherein X is G;
 - (iii) nucleotide 732 wherein X is C;
 - (iv) nucleotide 1293 wherein X is G;
 - (v) nucleotide 1322 wherein X is G;
 - (vi) nucleotide 1379 wherein X is C;

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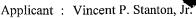
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(vii) nucleotide 1590 wherein X is T; (viii) nucleotide 1688 wherein X is G; (ixi) nucleotide 2401 wherein X is G; (x) nucleotide 2429 wherein X is A; (xi) nucleotide 2488 wherein X is T; (xii) nucleotide 2594 wherein X is T; (xiii) nucleotide 2618 wherein X is A; (xiv) nucleotide 3083 wherein X is A; (xv) nucleotide 3125 wherein X is A; (xvi) nucleotide 3212 wherein X is T (xvii) nucleotide 3619 wherein X is A; (xviii) nucleotide 3635 wherein X is A; (xix) nucleotide 4256 wherein X is A; (xx) nucleotide 4898 wherein X is G; (xxi) nucleotide 5006 wherein X is T; (xxii) nucleotide 5062 wherein X is A; (xxiii) nucleotide 5167 wherein X is A; (xxiv) nucleotide 11069 wherein X is G; (xxv) nucleotide 11238 wherein X is T; (xxvi) nucleotide 11293 wherein X is G; (xxvii) nucleotide 11422 wherein X is C; (xxviii) nucleotide 11686 wherein X is T; (xxix) nucleotide 12598 wherein X is C; (xxx) nucleotide 13171 wherein X is C, (xxxi) nucleotide 13298 Wherein X is A; (xxxii) nucleotide 13645 wherein X is C; (xxxiii) nucleotide 13751 wherein X is A; (xxxiv) nucleotide 13782 Wherein X is C; (xxxv) nucleotide 13806 wherein X is \mathcal{E} ; (xxxvi) nucleotide 13813 wherein X/Is C; (xxxvii) nucleotide 14479 wherein X is G; (xxxviii) T is inserted after nucleotide 14546; (xxxix) nucleotide 14585 wherein X is T; (xl) nucleotide 14729 wherein X is A; (xli) nucleotide 14787 wherein X is T; (xlii) nucleotide 14795 wherein X is A; (xliii) nucleotide 15041 wherein X is C; (xliv) nucleotide 15343 wherein X is A; (xlv) nucleotide 15449 wherein X is A; (xlvi) nucleotide 15502 wherein X is A; (xlvii) nucleotide 15545 wherein X is T; (xlvii) nucleotide 15589 wherein X is G; (xlix) nucleotide 15769 wherein X is T; (1) nucleotide 15839 wherein X is G;



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(li) nucleotide 16148 wherein X is A;

(lii) nucleotide 16198 wherein X is G; and

(liii) nucleotide 16202 wherein X is T

or the complement thereof; and

(c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.

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89./05 An isolated nucleic acid probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:9, the probe comprising at least one of:

- (a) nucleotide 128 wherein X is A
- (b) nucleotide 189 wherein X is G;
- (c) nucleotide 524 wherein X is G
- (d) nucleotide 1399 wherein X is A;
- (e) nucleotide 1464 wherein X is A
- (f) nucleotide 1636 wherein X is T
- (g) nucleotide 1738 wherein X is T and
- (h) nucleotide 2259 wherein X is C or the complement thereof.

96. 106 The isolated nucleic acid probe of claim 89 comprising at least two of:

- (a) nucleotide 128 wherein X is A;
- (b) nucleotide 189 wherein X is G;
- (c) nucleotide 524 wherein X is G;
- (d) nucleotide 1399 wherein X is A
- (e) nucleotide 1464 wherein X is ♠;
- (f) nucleotide 1636 wherein X is T
- (g) nucleotide 1738 wherein X is T and
- (h) nucleotide 2259 wherein X is &; or the complement thereof.

91./87 The probe of claim, 89 comprising no more than 500 contiguous nucleotides of SEQ ID NO:9.

92./08 The probe of claim 89 comprising no more than 200 contiguous nucleotides of SEQ ID NO:9.

93./09 The probe of claim 89 comprising no more than 100 contiguous nucleotides of SEQ ID NO:9.

94./10 The probe of claim 89 comprising no more than 50 contiguous nucleotides of SEQ 1D NO:9.

95.// The probe of claim 89 comprising DNA.

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96.//2The probe of claim 89 comprising a peptide nucleic acid.

97.//3 The probe of claim 89 further comprising a detectable label.

98.//4 The probe of claim 97 wherein the detectable label is a fluorescent label.

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99.//5 A method comprising:

(a) providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;

(b) contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:10, the probe comprising at least one of:

(i) nucleotide 128 wherein X is A;

- (ii) nucleotide 189 wherein X is G;
- (iii) nucleotide 524 wherein X is G;
- (iv) nucleotide 1399 wherein X is A;
- (v) nucleotide 1464 wherein X is A;
- (vi) nucleotide 1636 wherein X is T;
- (vii) nucleotide 1738 wherein X is T; and
- (viii) nucleotide 2259 wherein X is C;

or the complement thereof; and

(c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.

100. An isolated nucleic acid probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:10, the probe comprising at least one of:

- (a) nucleotide 183 wherein X is A;
- (b) nucleotide 483 wherein X is T;
- (c) nucleotide 601 wherein X is C; and
- (d) nucleotide 1299 wherein X is A; or the complement thereof.

101! The isolated nucleic acid probe of claim 100 comprising at least two of:

- (a) nucleotide 183 wherein X is A;
- (b) nucleotide 483 wherein X is T;
- (c) nucleotide 601 wherein X is C; and
- (d) nucleotide 1299 wherein X is A; or the complement thereof.

102. The probe of claim 100 comprising no more than 500 contiguous nucleotides of SEQ ID NO:10.

103.//9 The probe of claim 100 comprising no more than 200 contiguous nucleotides of SEQ ID NO:10.

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104. The probe of claim 100 comprising no more than 100 contiguous nucleotides of SEQ ID NO:10.

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103. The probe of claim 100 comprising no more than 50 contiguous nucleotides of SEO ID NO:10.

106. The probe of claim 100 comprising DNA.

107.123 The probe of claim 100 comprising a peptide nucleic acid.

108.124 The probe of claim 100 further comprising a detectable label.

109./25 The probe of claim 108 wherein the detectable label is a fluorescent label.

140./26A method comprising:

(a) providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;

(b) contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO: 0, the probe comprising at least one of:

(i) nucleotide 183 wherein X is A;

(ii) nucleotide 483 wherein X is I

(iii) nucleotide 601 wherein X is C;

(iv) nucleotide 1299 wherein X is A; or the complement thereof; and

(c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.

An isolated nucleic acid probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:11, the probe comprising at least one of:

(a) nucleotide 124 wherein X is T;

(b) nucleotide 439 wherein X is A;

(c) CT is inserted after nucleotide 1044;

(d) nucleotide 1331 wherein X is A;

(e) nucleotide 1977 wherein X is A;

(f) nucleotide 2149 wherein X is A;

(g) nucleotide 2467 wherein X is G;

(h) nucleotide 2634 wherein X is G;

(i) nucleotide 2975 wherein X is A;

(j) nucleotide 3116 wherein X is T;

(k) nucleotide 3255 wherein X is C;

(1) nucleotide 3344 wherein X is C;

(m) nucleotide 4051 wherein X is A; (n) nucleotide 4782 wherein X is A;

(o) nucleotide 5022 wherein X is C;

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- (p) nucleotide 5266 wherein X is A;
- (q) nucleotide 5285 wherein X is G;
- (r) nucleotide 5438 wherein X is A;
- (s) nucleotide 5482 wherein X is T;
- (t) nucleotide 5629 wherein X is A;
- (u) nucleotide 5648 wherein X is T; and
- (v) nucleotide 5731 wherein X is A; or the complement thereof.

112.128 The isolated nucleic acid probe of claim 111 comprising at least two of:

- (a) nucleotide 124 wherein X is T;
- (b) nucleotide 439 wherein X is A;
- (c) CT is inserted after nucleptide 1044;
- (d) nucleotide 1331 wherein X is A;
- (e) nucleotide 1977 wherein X is A;
- (f) nucleotide 2149 wherein X is A;
- (g) nucleotide 2467 wherein X is G
- (h) nucleotide 2634 wherein X is G;
- (i) nucleotide 2975 wherein XVs A;
- (j) nucleotide 3116 wherein X is T;
- (k) nucleotide 3255 wherein X is C;
- (1) nucleotide 3344 wherein X is C;
- (m) nucleotide 4051 wherein X is A;
- (n) nucleotide 4782 wherein X is A;
- (o) nucleotide 5022 wherein X is C
- (p) nucleotide 5266 wherein X is A;
- (q) nucleotide 5285 wherein X is G;
- (r) nucleotide 5438 wherein X is
- (s) nucleotide 5482 wherein X is
- (t) nucleotide 5629 wherein X is A
- (u) nucleotide 5648 wherein X is T; and
- (v) nucleotide 5731 wherein X is A; or the complement thereof.

113. 129 The probe of claim 117 comprising no more than 500 contiguous nucleotides of SEO ID NO:11.

114./30 The probe of claim 111 comprising no more than 200 contiguous nucleotides of SEO ID NO:11.

145.¹³ The probe of claim 111 comprising no more than 100 contiguous nucleotides of SEQ ID NO:11.

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The probe of claim 111 comprising no more than 50 contiguous nucleotides of SEQ ID NO:11.

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117. 13.3 The probe of claim 111 comprising DNA.

118.13 The probe of claim 111 comprising a peptide nucleic acid.

149./35The probe of claim 141 further comprising a detectable label.

120./36 The probe of claim 119 wherein the detectable label is a fluorescent label.

121./37A method comprising:

(a) providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;

- (b) contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:12, the probe comprising at least one of:
 - (i) nucleotide 124 wherein X is Γ;
 - (ii) nucleotide 439 wherein X is A
 - (iii) CT is inserted after nucleotide 1044;
 - (iv) nucleotide 1331 wherein X is A;
 - (v) nucleotide 1977 wherein X is A;
 - (vi) nucleotide 2149 wherein X is A
 - (vii) nucleotide 2467 wherein X is G;
 - (viii) nucleotide 2634 wherein X is G;
 - (ix) nucleotide 2975 wherein X is A;
 - (x) nucleotide 3116 wherein X is The
 - (xi) nucleotide 3255 wherein X is \$\dagger\$;
 - (xii) nucleotide 3344 wherein X is C
 - (xiii) nucleotide 4051 wherein X is A;
 - (xiv) nucleotide 4782 wherein X is A:
 - (xv) nucleotide 5022 wherein X is C;
 - (xvi) nucleotide 5266 wherein X is A;
 - (xvii) nucleotide 5285 wherein X is G
 - (xviii) nucleotide 5438 wherein X is A
 - (xix) nucleotide 5482 wherein X is T;
 - (xx) nucleotide 5629 wherein X is A;
 - (xxi) nucleotide 5648 wherein X is T; and
 - (xxii) nucleotide 5731 wherein X is A;

or the complement thereof; and

(c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.

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An isolated nucleic acid probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:12, the probe comprising at least one of:

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- (a) nucleotide 803 wherein X is T;
- (b) nucleotide 1747 wherein X is T; and
- (c) nucleotide 1900 wherein X is C; or the complement thereof.

123.¹³⁹The isolated nucleic acid probe of claim 122 comprising at least two of:

- (a) nucleotide 803 wherein X is T;
- (b) nucleotide 1747 wherein X is T; and
- (c) nucleotide 1900 wherein X is C; or the complement thereof.

124. The probe of claim 122 comprising no more than 500 contiguous nucleotides of SEQ ID NO:12.

125. The probe of claim 122 comprising no more than 200 contiguous nucleotides of SEQ ID NO:12.

126. The probe of claim 122 comprising no more than 100 contiguous nucleotides of SEQ ID NO:12.

127. The probe of claim 122 comprising no more than 50 contiguous nucleotides of SEQ ID NO:12.

128. The probe of claim 122 comprising DNA.

129. The probe of claim 122 comprising a peptide nucleic acid.

130. 146 The probe of claim 122 further comprising a detectable label.

131. The probe of claim 130 wherein the detectable label is a fluorescent label.

132.148 A method comprising:

- (a) providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;
- (b) contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:12, the probe comprising at least one of:
 - (a) nucleotide 803 wherein X is T;
 - (b) nucleotide 1747 wherein X is T; and
- (c) nucleotide 1900 wherein X is C; or the complement thereof; and

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(c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.

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An isolated nucleic acid probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:13, the probe comprising at least one of:

- (a) nucleotide 1424 wherein X is A;
- (b) nucleotide 1649 wherein X is A; and
- (c) nucleotide 2554 wherein X is G; or the complement thereof.

134. The isolated nucleic acid probe of claim 133 comprising at least two of:

- (a) nucleotide 1424 wherein X is A;
- (b) nucleotide 1649 wherein X is A; and
- (c) nucleotide 2554 wherein X is G; or the complement thereof.

135. The probe of claim 138 comprising no more than 500 contiguous nucleotides of SEQ ID NO:13.

136. The probe of claim 133 comprising no more than 200 contiguous nucleotides of SEQ ID NO:13.

137. The probe of claim 133 comprising no more than 100 contiguous nucleotides of SEQ ID NO:13.

138. The probe of claim 133 comprising no more than 50 contiguous nucleotides of SEQ ID NO:13.

139. The probe of claim 133 comprising DNA.

140. 156 The probe of claim 133 comprising a peptide nucleic acid.

141. 15 The probe of claim 133 further comprising a detectable label.

142. 156 The probe of claim 137 wherein the detectable label is a fluorescent label.

143./5% method comprising:

- (a) providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;
- (b) contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:13, the probe comprising at least one of:
 - (a) nucleotide 1424 wherein X is A;
 - (b) nucleotide 1649 wherein X is A; and
 - (c) nucleotide 2554 wherein X is G;

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or the complement thereof; and

(c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.

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144. An isolated nucleic acid probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:14, the probe comprising at least one of:

- (a) nucleotide 263 wherein X is G;
- (b) nucleotide 266 wherein X is T;
- (c) nucleotide 527 wherein X is G;
- (d) nucleotide 1037 wherein X is G;
- (e) nucleotide 1139 wherein X is A;
- (f) nucleotide 1217 wherein X is T;
- (g) nucleotide 1647 wherein X is T;
- (h) nucleotide 1955 wherein X is A;
- (i) nucleotide 2017 wherein X is A;
- (j) nucleotide 2037 wherein X is A;
- (k) nucleotide 2189 wherein X is G;
- (1) nucleotide 2282 wherein X is T; and
- (m) nucleotide 2309 wherein X is G or the complement thereof.

The isolated nucleig acid probe of claim 144 comprising at least two of:

- (a) nucleotide 263 whetein X is G;
- (b) nucleotide 266 wherein X is T;
- (c) nucleotide 527 wherein X is G:
- (d) nucleotide 1037 wherein X is G;
- (e) nucleotide 1139 wherein X is A;
- (f) nucleotide 1217 wherein X is T;
- (g) nucleotide 1647 wherein \mathbf{X} is T;
- (h) nucleotide 1955 wherein X is A;
- (i) nucleotide 2017 wherein X is A;
- (j) nucleotide 2037 wherein X\is A;
- (k) nucleotide 2189 wherein X is G;
- (1) nucleotide 2282 wherein X is T; and

(m) nucleotide 2309 wherein X is G; or the complement thereof.

146. The probe of claim 144 domprising no more than 500 contiguous nucleotides of SEQ ID NO:14.

147. The probe of claim 144 comprising no more than 200 contiguous nucleotides of SEQ ID NO:14.

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148. The probe of claim 144 comprising no more than 100 contiguous nucleotides of SEQ ID NO:14.

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149. The probe of claim, 144 comprising no more than 50 contiguous nucleotides of SEQ ID NO:14.

150. The probe claim 144 comprising DNA.

The probe of claim 144 comprising a peptide nucleic acid.

152. The probe of claim 144 further comprising a detectable label.

-153!/69 The probe of claim 1.52 wherein the detectable label is a fluorescent label.

154/10 A method comprising

- (a) providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;
- (b) contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:14, the probe comprising at least one of:
 - (i) nucleotide 263 wherein X is G;
 - (ii) nucleotide 266 wherein X is T;
 - (iii) nucleotide 527 wherein X is G;
 - (iv) nucleotide 1037 wherein X is G;
 - (v) nucleotide 1139 wherein X is A;
 - (vi) nucleotide 1217 wherein X is T;
 - (vii) nucleotide 1647 wherein X is T;
 - (viii) nucleotide 1955 wherein X is A; (ix) nucleotide 2017 wherein X is A;
 - (x) nucleotide 2037 wherein X is A:
 - (xi) nucleotide 2189 wherein X is G;
 - (xii) nucleotide 2282 wherein X is T; and
 - (xiii) nucleotide 2309 wherein X is G;

or the complement thereof; and

(c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.

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An isolated nucleic acid probe comprising at least 15 contiguous nucleotides of he nucleotide sequence of SEQ ID NO:15, the probe comprising at least one of:

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- (a) nucleotide 120 wherein X is C;
- (b) nucleotide 464 wherein X is G;
- (c) nucleotide 519 wherein X is T;
- (d) nucleotide 668 wherein X is T;
- (e) nucleotide 1059 wherein X is C;
- (f) nucleotide 1289 wherein X is A;
- (g) nucleotide 1308 wherein X is C;
- (h) nucleotide 1784 wherein X is A;

or the complement thereof.

The isolated nucleic acid probe of claim 155 comprising at least two of:

- (a) nucleotide 120 wherein X is C;
- (b) nucleotide 464 wherein X is G;
- (c) nucleotide 519 wherein X is T;
- (d) nucleotide 668 wherein X is T;
- (e) nucleotide 1059 wherein X is C;
- (f) nucleotide 1289 wherein X is A;
- (g) nucleotide 1308 wherein X is C;
- (h) nucleotide 1784 wherein X is A;

or the complement thereof.

173 171 The probe of claim 155 comprising no more than 500 contiguous nucleotides of SEQ ID NO:15.

158. The probe of claim 155 comprising no more than 200 contiguous nucleotides of SEQ ID NO:15.

The probe of claim 155 comprising no more than 100 contiguous nucleotides of SEO ID NO:15.

160. The probe of claim 155 comprising no more than 50 contiguous nucleotides of SEQ ID NO:15.

161. 17 The probe claim 155 comprising DNA.

162. 79 The probe of claim 155 comprising a peptide nucleic acid.

163.180 The probe of claim 155 further comprising a detectable label.

164. 181 The probe of claim 163 wherein the detectable label is a fluorescent label.

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-165. A method comprising:

(a) providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;

(b) contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQID NO:15, the probe comprising at least one of:

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- (i) nucleotide 120 wherein X is C;
- (ii) nucleotide 464 wherein X is G:
- (iii) nucleotide 519 wherein X is T;
- (iv) nucleotide 668 wherein X is T;
- (v) nucleotide 1059 wherein X is C;
- (vi) nucleotide 1289 wherein X is A;
- (vii) nucleotide 1308 wherein X is C;
- (vii) nucleotide 1784 wherein X is A;

or the complement thereof; and

(c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.

An isolated nucleic acid probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQUD NO:16, the probe comprising at least one of:

- (a) nucleotide 575 wherein X is C;
- (b) nucleotide 648 wherein X is C;
- (c) nucleotide 771 wherein X is C;
- (d) nucleotide 883 wherein X is A;
- (e) C is inserted after nucleotide 941; and
- (f) nucleotide 1051 wherein X is C; or the complement thereof.

167. The isolated nucleic acid probe of claim 166 comprising at least two of:

- (a) nucleotide 575 wherein X is C
- (b) nucleotide 648 wherein X is C
- (c) nucleotide 771 wherein X is C
- (d) nucleotide 883 wherein X is A
- (e) C is inserted after nucleotide 941; and
- (f) nucleotide 1051 wherein X is C

or the complement thereof.

168. The probe of claim 166 comprising no more than 500 contiguous nucleotides of SEQ ID NO:16.

The probe of claim 166 comprising no more than 200 contiguous nucleotides of SEQ ID NO:16.

170. The probe of claim 166 comprising no more than 100 contiguous nucleotides of SEQ ID NO:16.

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171. The probe of claim 166 comprising no more than 50 contiguous nucleotides of SEQ ID NO:16.

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172. The probe of claim 166 comprising DNA.

173. The probe of claim 166 comprising a peptide nucleic acid.

174. The probe of claim 166 further comprising a detectable label.

175.192 The probe of claim 174 wherein the detectable label is a fluorescent label.

-176. ¹⁹³ A method comprising:

- (a) providing a sample comprising nucleic acid molecules present in a biological sample obtained from a patient;
- (b) contacting the sample with a probe comprising at least 15 contiguous nucleotides of the nucleotide sequence of SEQ ID NO:16, the probe comprising at least one of:
 - (i) nucleotide 575 wherein X is C;
 - (ii) nucleotide 648 wherein X is C;
 - (iii) nucleotide 771 wherein X is C;
 - (iv) nucleotide 883 wherein X is A;
 - (v) C is inserted after nucleotide 941; and
- (vi) nucleotide 1051 wherein X is C; or the complement thereof; and
- (c) determining if the sample comprises a nucleic acid molecule that hybridizes to the probe.--

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